

Ms. Bibb 54, 7000014110

Query Match	2.2%	5.7%	32.6%	GP 4	2.4%	1.00%
Best Local Similarity	56.0%			Pred No 0.08%		
Matches	51	Conservative	10	MisMatches	10	Indels
						Gaps
						0





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[illegible]

RESULT 9  
US 04-905-223-137  
Sequence 137, Application US/08905223  
Patent No. 6126929  
GENERAL INFORMATION:  
APPLICANT: Edwards, Jean Baptiste D.  
APPLICANT: Duclouet, Aymeric  
APPLICANT: Lacroix, Bruno  
TITLE OF INVENTION: 5' ESTS FOR SECRETED PROTEINS  
NUMBER OF SEQUENCES: 503  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: Kachle, Marten, Simon & Boar  
STREET: 501 West Broadway  
CITY: San Diego  
STATE: California  
COUNTRY: USA  
ZIP: 92101-3505  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: Win95  
SOFTWARE: Word  
INVENTOR AFFILIATION DATA:  
AFFILIATION NUMBER: 03/02/001.223  
FILING DATE:  
CLASSIFICATION: 516  
ATTORNEY/AGENT INFORMATION:  
NAME: Israelien, Ned A.  
REGISTRATION NUMBER: 29,655  
REFERENCE/DOC# NUMBER:



Query Matrix	7.8	50.10	1.1	34.7	100.00
Best Local Similarity	47.08				
Pred. No.	5.8				

[illegible]

[illegible]

RESULT 15  
 US-09-153-599A-2  
 ; Sequence 2, Application US/09153599A  
 ; Patent No. 6420177  
 ; GENERAL INFORMATION:  
 ; APPLICANT: Weber, J. Mark  
 ; APPLICANT: Lau, B. Minh  
 ; TITLE OF INVENTION: Method for Strain Improvement of  
 ; TITLE OF INVENTION: Extracellular Production Protein

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RESULT 14
US-08-366-276-1/c
Sequence 1, Application US/0936C276
Patent No. 5534409
GENERAL INFORMATION:
APPLICANT: Groner, Bernd
APPLICANT: Gouilleux, Fabrice
APPLICANT: Wakao, Hiroshi
TITLE OF INVENTION: Cytokine Regulated Transcription Factor
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: CIAA-GELBY Corporation
STREET: 7 Skyline Drive
City: Hawthorne
STATE: New York
COUNTRY: USA
ZIP: 10532.
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.30B
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/366,276
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: GB 9409396.0
FILING DATE: 11-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: Elmer, James Scoot
REGISTRATION NUMBER: 36,129
REFERENCE/DOCKET NUMBER: 4-19992/A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (919) 541-8614
TELEFAX: (919) 541-8689
INFORMATION FOR SEQ ID NO.: 1.
SEQUENCE CHARACTERISTICS:
LENGTH: 2818 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
FEATURES:
NAME/KEY: CDS
LOCATION: 244..2625
OTHER INFORMATION: /product= "mammary gland factor"
US-08-366-276-1

Query Match           6.9%; Score 33.6; DB 1; Length 2818;
Best Local Similarity   53.4%; Pred.No. 3.4;
Matches    63; Conservative      0; Mismatches     55; Indels       0; Gaps        0,

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CORRESPONDENCE ADDRESS:
ADDRESSEE: Pockey, Milnamow & Katz, Ltd.
STREET: 180 N. Streteon Avenue, 2 Prudentia Plaza
CITY: Chicago
STATE: Illinois
COUNTRY: U.S.A.
ZIP: 60601

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Parent'n Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/153,599A
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Mueller, Lisa V.
REGISTRATION NUMBER: 38,978
REFERENCE/DOCKET NUMBER: EPR2159D004IUS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-616-5400
TELEFAX: 312-616-5460
INFORMATION FOR SEQ ID NO.: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1206 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
FEATURE:
NAME/KEY: CDS
LOCATION: 1..1203
US-09-153-599A-2

Query Match: 6.9%; Score 33.2; DB 4; Length 1206;
Best Local Similarity 48.2%; Pos No 3 5;
Matches 92; Conservative 0; Mismatches 99; Indels 0; Gaps 0;

CY 178 ATCTGCGGGGGTGTGAGTCAGACCGGCGGGGGTGTGCGGGTTGTTTGTTTGGTGGG 237
DB 572 AGGTTGAAGGGGAGGGATGAAAGGGGGGGGGTGTGCGAGGGGGTGTGATTAATGAGGGTGGCAACGTGG 631
CY 238 AGAGCTTGAAGTAGGTGTGTGTGCGGGGGTGTGTAAGAAGCGGGGGTATTTTGTGAGAGGGCT 297
DB 632 AGCTGGGAGAGATGGTACAGAGTGGGGGGTGTGTGTATACAGCGGGGGTATTTGGTGAATA 691
CY 298 GGAAGCGGTTGGTGGTGTGATGTGATGTGAGAACAGGGGGTGTGTTTGGTGTGGGGTGGAGATAG 357
DB 692 TGGCGGAGTTGGTGTGTGTGAGAGCATGGCGAGAGAGTATCTGGGGGTGATGAGCAAGGTGG 751
CY 358 AGGCCACGGTC 368
DB 752 TCGCCACGGC 762

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